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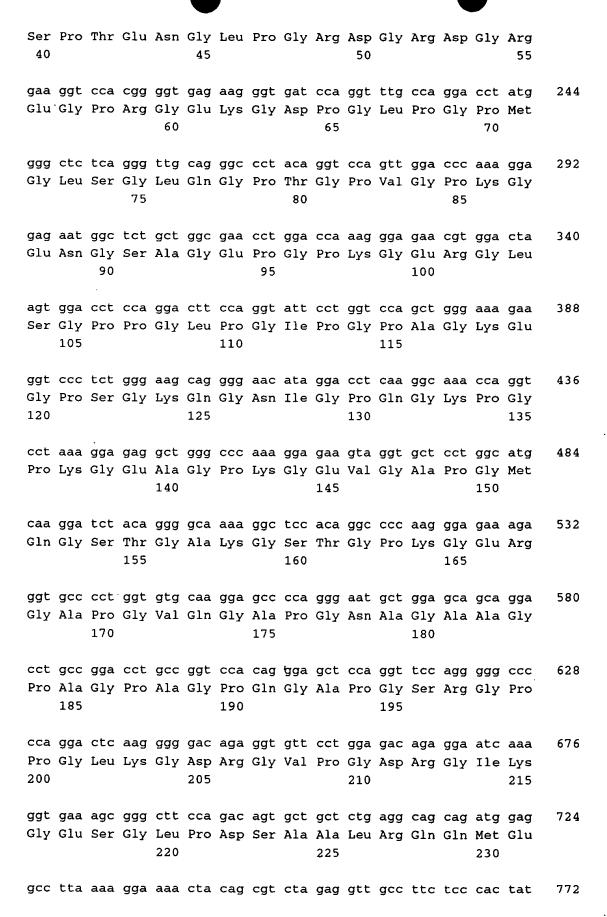
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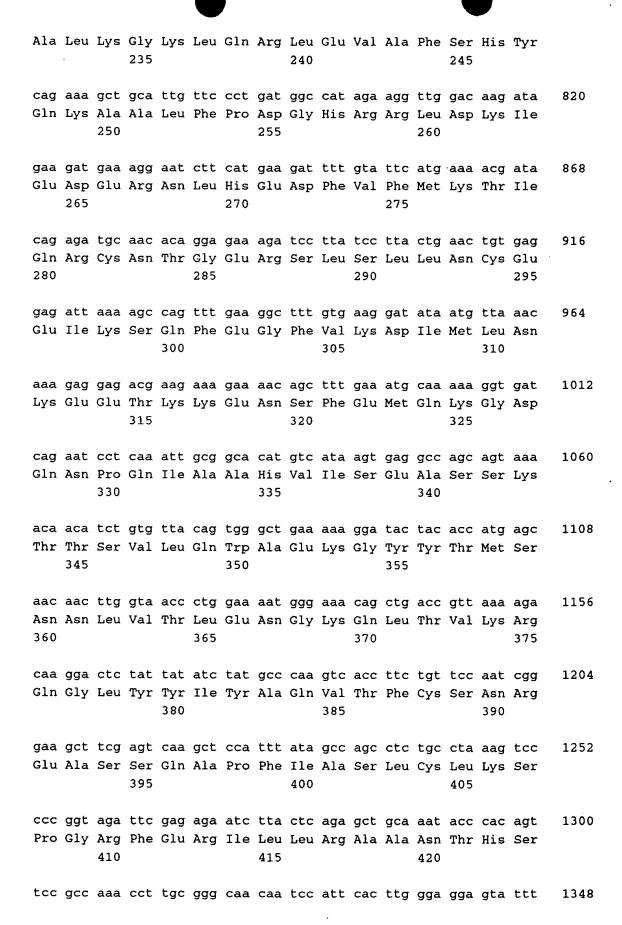
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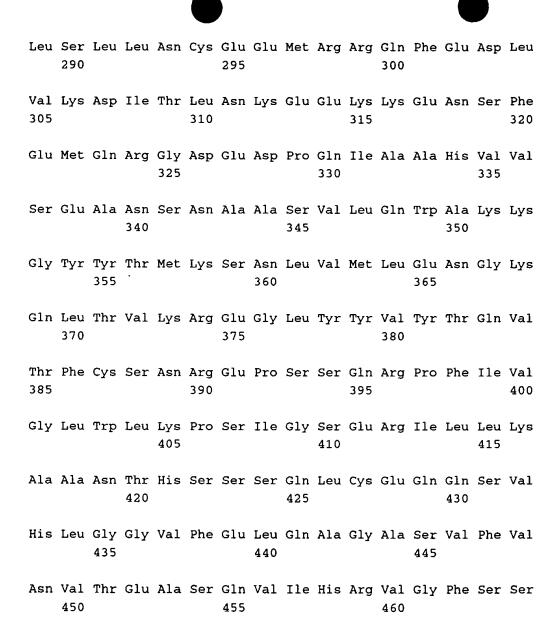
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<302> Recombinant human CD40 ligand stimulates B cell
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<303> J. Exp. Med.
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<223> Description of Artificial Sequence: Murine surfactant protein D (without the CRD) fused to the extracellular portion of human CD40L

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275

265

280

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285

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Leu Ser Leu Leu Asn Cys Glu Glu Ile Lys Ser Gln Phe Glu Gly Phe 295 300 Val Lys Asp Ile Met Leu Asn Lys Glu Glu Thr Lys Lys Glu Asn Ser 310 315 Phe Glu Met Gln Lys Gly Asp Gln Asn Pro Gln Ile Ala Ala His Val 325 330 335 Ile Ser Glu Ala Ser Ser Lys Thr Thr Ser Val Leu Gln Trp Ala Glu 345 350 Lys Gly Tyr Tyr Thr Met Ser Asn Asn Leu Val Thr Leu Glu Asn Gly 360 Lys Gln Leu Thr Val Lys Arg Gln Gly Leu Tyr Tyr Ile Tyr Ala Gln 370 375 380 Val Thr Phe Cys Ser Asn Arg Glu Ala Ser Ser Gln Ala Pro Phe Ile 390 395 400 Ala Ser Leu Cys Leu Lys Ser Pro Gly Arg Phe Glu Arg Ile Leu Leu 410 Arg Ala Ala Asn Thr His Ser Ser Ala Lys Pro Cys Gly Gln Gln Ser 420 425 430 Ile His Leu Gly Gly Val Phe Glu Leu Gln Pro Gly Ala Ser Val Phe 435 440 445 Val Asn Val Thr Asp Pro Ser Gln Val Ser His Gly Thr Gly Phe Thr 455 460 450 Ser Phe Gly Leu Leu Lys Leu 470 465 <210>3 <211> 1574 1 <212> DNA <213> Artificial Sequence <220> <221> 5'UTR <222>(7)..(31) <223> 5' UTR taken from rat sequence for surfactant

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protein D

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   Guo, N
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   14.
<303> J. Immunol.
<304> 155
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   Dougall, WC.
<302> A homologue of the TNF receptor and its ligand enhance
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gcc tta aaa gga aaa cta cag cgt cta gag gtt gcc ttc tcc cac tat 772
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235 240 245

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Arg Ile Ser Glu Asp Ser Thr His Cys Phe Tyr Arg Ile Leu Arg Leu
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130

145

135

150

140

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160

155

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<301> Motwani, M

Ser Gly Asn Ser Glu Phe His Phe Tyr Ser Ile Asn Val Gly Gly Phe 455 450 460 Phe Lys Leu Arg Ala Gly Glu Glu Ile Ser Ile Gln Val Ser Asn Pro 470 475 480 Ser Leu Leu Asp Pro Asp Gln Asp Ala Thr Tyr Phe Gly Ala Phe Lys 485 490 495 Val Gln Asp Ile Asp 500 <210>5 <211> 1477 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Murine surfactant protein D (except CRD) fused to the extracellular domain of murine CD40 ligand <220> <221> 5'UTR <222>(7)..(31) <223> 5' UTR from rat surfactant protein D <220> <221> sig_peptide <222> (32)..(88) <223> Signal peptide from murine surfactant protein D <220> <221> CDS <222> (32)..(1441) <220> <221> misc_recomb <222> (88)..(799) <223> Mature murine surfactant protein D including hub region, collagenous portion, and neck, but excluding carbohydrate recognition domain (CRD) <220> <221> misc_feature <222> (800)..(1441) <223> Murine CD40 ligand extracellular region, including stalk <300>

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   Guo, N
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<302> Mouse surfactant protein-D. cDNA cloning,
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<303> J. Immunol.
<304> 155
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   Clifford, K N.
<302> Molecular and biological characterization of a murine
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                  15
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80

85

75

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1477

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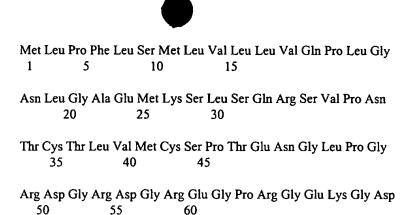
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<213> Artificial Sequence

<223> Description of Artificial Sequence: Murine surfactant protein D (except CRD) fused to the extracellular domain of murine CD40 ligand



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Pro Lys Gly Glu Arg Gly Leu Ser Gly Pro Pro Gly Leu Pro Gly Ile 100 105 110

Pro Gly Pro Ala Gly Lys Glu Gly Pro Ser Gly Lys Gln Gly Asn Ile 115 120 125

Gly Pro Gln Gly Lys Pro Gly Pro Lys Gly Glu Ala Gly Pro Lys Gly 130 135 140

Glu Val Gly Ala Pro Gly Met Gln Gly Ser Thr Gly Ala Lys Gly Ser 145 150 155 160

Thr Gly Pro Lys Gly Glu Arg Gly Ala Pro Gly Val Gln Gly Ala Pro . 165 170 175

Gly Asn Ala Gly Ala Ala Gly Pro Ala Gly Pro Ala Gly Pro Gln Gly 180 185 190

Ala Pro Gly Ser Arg Gly Pro Pro Gly Leu Lys Gly Asp Arg Gly Val 195 200 205

Pro Gly Asp Arg Gly Ile Lys Gly Glu Ser Gly Leu Pro Asp Ser Ala 210 215 220

Ala Leu Arg Gln Gln Met Glu Ala Leu Lys Gly Lys Leu Gln Arg Leu 225 230 235 240

Glu Val Ala Phe Ser His Tyr Gln Lys Ala Ala Leu Phe Pro Asp Gly 245 250 255

His Arg Arg Leu Asp Lys Val Glu Glu Val Asn Leu His Glu Asp 260 265 270

Phe Val Phe Ile Lys Lys Leu Lys Arg Cys Asn Lys Gly Glu Gly Ser 275 280 285



Leu Ser Leu Leu Asn Cys Glu Glu Met Arg Arg Gln Phe Glu Asp Leu 290 295 300

Val Lys Asp Ile Thr Leu Asn Lys Glu Glu Lys Lys Glu Asn Ser Phe 305 310 315 320

Glu Met Gln Arg Gly Asp Glu Asp Pro Gln Ile Ala Ala His Val Val 325 330 335

Ser Glu Ala Asn Ser Asn Ala Ala Ser Val Leu Gln Trp Ala Lys Lys 340 345 350

Gly Tyr Tyr Thr Met Lys Ser Asn Leu Val Met Leu Glu Asn Gly Lys 355 360 365

Gln Leu Thr Val Lys Arg Glu Gly Leu Tyr Tyr Val Tyr Thr Gln Val

Thr Phe Cys Ser Asn Arg Glu Pro Ser Ser Gln Arg Pro Phe Ile Val 385 390 395 400

Gly Leu Trp Leu Lys Pro Ser Ile Gly Ser Glu Arg Ile Leu Leu Lys 405 410 415

Ala Ala Asn Thr His Ser Ser Ser Gln Leu Cys Glu Gln Gln Ser Val 420 425 430

His Leu Gly Gly Val Phe Glu Leu Gln Ala Gly Ala Ser Val Phe Val 435 440 445

Asn Val Thr Glu Ala Ser Gln Val Ile His Arg Val Gly Phe Ser Ser 450 455 460

Phe Gly Leu Leu Lys Leu 465 470